

Amendments to the Specification

Please replace Table 1 on page 13 of the specification with the following amended

Table 1:

Functional elements in the RSus3 promoter region	Position	Position (-intron)	RSus3 sequence ¹	Consensus sequence	Reference (Consensus sequence)
Translation start site	+1	+1	CAATGG (SEQ ID No 7)	CAATGG (SEQ ID No 8)	[Joshi 1987]
Intron 1 (Acceptor splice site)	-27		TCCAG GC (SEQ ID No 9)	TGCAG GT (SEQ ID No 10)	[Simpson & Filipowicz 1996], consensus for monocots
Intron 1 (Donor splice site)	-892		AG GTAGAG (SEQ ID No 11)	AG GTAAGT (SEQ ID No 12)	
TATA box	-986	-121	TATAAATA (SEQ ID No 13)	TATATATA (SEQ ID No 14)	[Joshi 1987]
CAAT box	-999	-134	GCACATTT (SEQ ID No 15)	GGNCAATC (SEQ ID No 16)	
GCN4 box	-1032	-167	GTGAGGC AG (SEQ ID No 17)	(G/A)TGA(G/C)TCA(T/G) (SEQ ID No 18)	[Muller & Knudsen 1993], cis-element involved in endosperm specificity
Endosperm boxes	-1072	-207	AGTAAAG (SEQ ID No 19)	TG(T/C/A)AA(G/A) (SEQ ID No 20)	[Marzábal et al. 1998], cis-element involved in endosperm specificity
	-1130	-265	TGCAAAC (SEQ ID No 21)		
	-1349	-484	TGTCAAA (SEQ ID No 22)		

Legumin boxes (RY repeats)	-1539	-674	CGTGCAT G (<u>SEQ ID No 23</u>)	CATGCATG (<u>SEQ ID No 24</u>)	[Bäumlein et al. 1992] cis-element involved in seed specificity
	-1586	-721	CATGTATG (<u>SEQ ID No 25</u>)		
	-1624	-759	CATGCAT A (<u>SEQ ID No 26</u>)		
	-1707	-842	CATGCAT GCATG (<u>SEQ ID No 27</u>)		
	-2115	-1250	CATGCCTG (<u>SEQ ID No 28</u>)		
	-2480	-1615	CAGGCAT GCATC (<u>SEQ ID No 29</u>)		